

SPINOCEREBELLAR ATAXIA TYPE 8 AND METHODS OF DETECTION

ABSTRACT OF THE DISCLOSURE

The present invention provides an isolated nucleic acid molecule containing a repeat region of an isolated spinocerebellar ataxia type 8 (SCA8) coding sequence, the coding sequence located within the long arm of chromosome 13, and the complement of the nucleic acid molecule. Diagnostic methods based on identification of this repeat region are also provided.

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